





TECH CINIER 1800/2900

ClustalW Formatted Alignments

	SEQ SEQ SEQ SEQ	ID ID ID	2 3 4	M M M	L G A	L P S	L G P	L A R	L P S	L F S	A A G	P R Q	L V P	F G G	A L W P P	R P X	P L P	P P P	G L P	A L P	G V P	G V P	A M P	Q A A	T A R	P G L	N V L	A A L
	SEQ SEQ SEQ SEQ SEQ	ID ID ID	2 3 4	T P L	S V L	E W L	G A L	C S P	Q H L	I S L	I P L	H H P	P L L	P P A	L W R P	E P G	G H A	G S W	I R G	R V W	Y P A	R P R	G H G	L P A	T S P	R S R	D E P	Q R P
	SEC SEC SEC SEC SEC	ID ID ID	2 3 4	V R P	K A S	A V S	I Y P	N I P	F G L	L A S	P L I	V F M	D P G	Y M L	R E S M V	I G P	E G L	Y W T	V P K	C G E	R G V	G Q A	E A K	R C G	E Q S	Y P I	V A G	G V R
	SEG SEG SEG SEG	ID ID ID	2 3 4	P E G	K M V	V A L	R L P	K E A	C D V	L V E	A N L	N S A	G R I	S R E	LWDQN	T I I	D L R	M P N	D D E	T Y S	P E L	S L L	R K R	C L P	V I Y	R H F	I H L	םםי
	SEQ SEQ SEQ SEQ SEQ	ID ID ID	2 3 4	S S L	K K R	C S	Y D Y	L P D	T G T	L Q E	E A C	N T D	G K N	K Y A	L V L K A	F Y G	L E L	T L K	G L A	G Y F	D N Y	L D D	P P A	A I I	L K K	Q I Y	G I G	A L P
	SEQ SEQ SEQ SEQ SEQ	מו מו מו	2 3 4	R M N	V P H	D G L	F C M	R S V	C S F	D V G	P S G	D T V	F L C	H V P	S L A S D	V E V	G A T	S A S	S R J	R M I	\$ W A	I N E	C	S I L	Q > 0	G L G	Q S ₩	W Y N
•	SEQ SEQ SEQ SEQ	ID ID ID	2 3 4	S G L	T S V	PSQ	K S L	P P S	H A F	C L A	Q S A	V И Т	N R T	R Q P	I T R V	P F L	H P A	STD	E F K	R F K	R R K	A T Y	V H P	Y P Y	I S F	G A F	A T R	LLT

```
SEQID 1 NKNQFKSFLRTIPNDEHQATAMADI
SEQ ID 2
      F P M S G G W P G G Q A C Q P A V E M A L E D V N
      HNPTRVKLFEKWGWKKIATIQQTTE
SEQ ID 3
SEQ ID 4
      V P S D N A V N P A I L K L L K H Y Q W K R V G T
SEQID 5 YASTAPELS DNTRYDFFSRVVPPDS
SEQ ID 1
      IEYFRWNWVGTIAADDDYGRPGIEK
SEQ ID 2
      S R R D I L P D Y E L K L I H H D S K C D P G Q A
SEQ ID 3
      V F T S T L D D L E E R V K E A G I E I T F R Q S
SEQID 4 LTQDVQRFSEVRNDLTGVLYGEDIE
SEQID 5 YQAQAMVDIVTALGWNYVSTLASEG
SEQ ID 1
      FREEAEERDICIDFSELISQYSDEE
SEQ ID 2
      TKYLYELLYNDPIKIILMPGCSSVS
SEQ ID 3
      F F S D P A V P V K N L K R Q D A R I I V G L F Y
SEQ ID 4
      ISDTESFSNDPCTSVKKLKGNDVRI
SEQID 5 NYGESGVEAFTQISREIGGVCIAQS
SEQID 1 EIQH V V E V I Q N S T A K V I V V F S S G P D
SEQ ID 2
      TLVAEAARMWNLIVLSYGSSSPALS
      ETEARKVFCEVYKERLFGKKYVWFL
SEQ ID 3
SEQ ID 4 ILGQFDQNMAAKVFCCAYEENMYGS
SEQID 5 QKIPREPRPGEFEKIIKRLLETPNA
SEQ ID 1
      LEPLIKEIVRRNITGKIWLASEAWA
SEQID 2 NRQRFPTFFRTHPSATLHNPTRVKL
      IGWYADNWFKIYDPSINCTVDEMTE
SEQ ID 3
SEQID 4 KYQWIIPGWYEPSWWEQVHTEANSS
SEQID 5 RAVIMFANEDDIRRILEAAKKLNQS
SEQID 1 SSSLIAMPQYFHVVGGTIGFALKAG
SEQID 2 FEKWGWKKIATIQQTTEVFTSTLDD
SEQID 3 A V E G H I T T E I V M L N P A N T R S I S N M T
SEQID 4 RCLRKNLLAAMEGYIGVDFEPLSSK
SEQ ID 5 G H F L W I G S D S W G S K I A P V Y Q Q E E I A
SEQ ID 1
      QIPGFREFLKKVHPRKSVHNGFAKE
SEQ ID 2
      LEERVKEAGIEITFRQSFFSDPAVP
SEQ ID 3
      SQEFVEKLTKRLKRHPEETGGFQEA
SEQID 4 QIKTISGKTPQQYEREYNNKRSGVG
SEQID 5 EGAVTILPKRASIDGFDRYFRSRTL
```

TITLE: G-Protein Fusion Receptors And Chimeric Gaba, Receptors INVENTORS: Thomas STORMANN, et al. SN: 09/679,664 ATTV #: (072827-1801)

```
SEQID 1 FWEETFNCHLQEGAKGPLPVDTFLR
SEQ ID 2
      V K N L K R Q D A R I I V G L F Y E T E A R K V F
SEQ ID 3
      PLAYDAIWALALALNKTSGGGGRSG
SEQID 4 PSKFHGYAYDGIWVIAKTLQRAMET
SEQID 5 ANNRRNVWFAEFWEENFGCKLGSHG
SEQID 1 GHEESGDRFSNSSTAFRPLCTGDEN
SEQID 2 CEVYKERLFGKKYVWFLIGWYADNW
SEQID 3 V R L E D F N Y N N Q T I T D Q I Y R A M N S S S
SEQID 4 LHAS SRHQRIQD FNYTDHTLGRIIL
SEQID 5 KRNSHIKKCTGLERIARDSSYEQEG
SEQ ID 1
      ISSVETPYIDYTHLRISYNVYLAVY
SEQID 2 FKIYDPSINCTVDEMTEAVEGHITT
SEQID 3 FEGVSGHVVFDASGSRMAWTLIEQL
SEQID 4 NAMNETNFFGVTGQVVFRNGERMGT
SEQID 5 K V Q F V I D A V Y S M A Y A L H N M H K D L C P
SEQID 1 SIAHALQDIYTCLPGRGLFTNGSCA
SEQID 2 EIVMLNPANTRS I SNMTSQEFVEKL
SEQID 3 QGG S Y K K I G Y Y D S T K D D L S W S K T D K
SEQID 4 IKFTQFQDSREVKVGEYNAVADTLE
SEQID 5 GYIGL CPRMSTIDGKELLGYIRAVN
SEQID 1 DIKKVEAWQVLKHLRHLNFTNNMGE
SEQID 2 TKRLKRHPEETGGFQEAPLAYDAIW
SEQID3 WIGGSPPADQTLVIKTFRFLSQK
SEQ ID 4
      IINDTIRFQGSEPPKDKTIILEQLR
SEQID 5 FNGSAGTPVTFNENGDAPGRYDIFQ
SEQID 1 QVTFDECGDLVGNYSIINWHLSPED
SEQID 2 ALALALNKTS GGGGRSGVRLEDFNY
SEQ ID 3
SEQID4 KISLP
SEQ ID 5 YQITNKSTEYKVIGHWTNQLHLKVE
SEQID 1 GSIVFKEVGYYNVYAKKGERLFINE
SEQID 2 NNQTITDQIYRAMNSSSFEGVSGHV
SEQ ID 3
SEQ ID 4
SEQID 5 DMQWAHREHTHPASVCSLPCKPGER
```

SEQ ID 1 SEQ ID 2 SEQ ID 3	E V																								
SEQ ID 4 SEQ ID 5	ĸ	K	т	٧	ĸ	G	ν	P	С	С	w	н	С	E	R	С	E	G	Y	N	Y	Q	v	D	E
SEQ ID 1 SEQ ID 2 SEQ ID 3																									
SEQ ID 4 SEQ ID 5	L	s	С	Е	L	С	P	L	D	Q	R	P	N	М	N	R	Τ	G	С	Q	L	I	P	I	I
SEQ ID 1 SEQ ID 2 SEQ ID 3 SEQ ID 4														F Q		S	N	Ε	N	н	T	S	С	I	A
SEQ ID 5	K	L	E	W	Н	S	P	W																	
SEQ ID 1 SEQ ID 2 SEQ ID 3 SEQ ID 4 SEQ ID 5	К	E	I	Ε	F	L	S	w	Т	E	P	F													

THLE: G-Protein Fusion Receptors And Chimeric Gaba, Receptors INVENTORS: Thomas STORMANN, et al SN: 09/679,664 ATTY #: (072827-1801)

```
SE0 (C 6
       G LAUTUFAVUG LEUTAFVUG VELKERNTE L
       LF ISVSVLSSLG IVLAVVCLS FN I YNSHVR
SECIDIZ
       L F I S V S V L S S L G T V L A V V C L S F N T Y N S H V A
SEC ID 8
       LYSTISALTILGMIMASAFLFFNIKNANOK
5EQ ID 9
       A V V P V F V A I L G I I A T T F V I V T F V R Y N D T P I
SEQ 10-10
       V K A T N R E L S Y L L L F S L L C C F S S S L F F I G E P
SEQ ID 6
       YIONSOPHLNNLTAVGCSLALAAVFPLGLO
SEQID 7
SEQ ID 8
       Y I Q N S C P N L N N L T A V G C S L A L A A V F P L G L D
       LIKMSSPYMNNLIILGGMLSYASIFLFGLD
SEQ 10 9
      VRASGRELSYVLLTG I FLC YS I T FLM I A A P
SEQ 10 10
       Q D W T C R L R Q P A F G I S F V L C I S C I L V K T N R V
SEQ 10 6
SEQ ID 7
       GYHIGRNOFPFVCOARLWLLGLGFSLGYG5
       G Y H I G R N O F P F V C Q A R L W L L G L G F S L G Y G S
SEQ ID 8
       GSFVSEKTFETLCTVRTWILTVGYTTAFGA
SEQ ID 9
SEQID 10 DTIICS FRRV FLG LGMC FS YAALLTK TNR I
       LLVFEAKIPTSFHRKWWGLNLQFLLVFLCT
SEO ID 6
       M F T K I W W V H T V F T K K E E K K E W R K T L E P W K L
SEQ ID 7
       MFTKIWWVHTVFTKKEEKKEWAKTLEPWKL
SEQ ID 8
       MFAKTWRVHAIFKNVKMKKKIIKOQKELVI
SEO ID 9
SEOID 10 HRIFE OGKKSVTAPKFISPASOLVITFSLI
       FMQ LV 1 CV I W L Y TAPPS S Y R N Q E L E D E I I F
SEQ 10 6
SEQ ID 7
       YATVGLLVGMDVLTLAIWQIVDPLHRTIET
       YATVGLLVGMDVLTLAIWOIVDPLHATIE
550 ID 8
SEQ ID 9
       V G G M L L I D L C I L I C W Q A V O P L R R T V E K Y S M
SEOID 10 S V O L L G V F V W F V V D P P H I I I D Y G E Q R T L O P
       ITCHEGSLMALGFLIGYTCLLAAICFFFAF
SEQ ID 6
       FAKEEPKEO I O V S I L P G L E H C S S R K M N T W L
SEQ ID 7
       FAKEEPKED LOVS I LPGLEHCSSAKMNTW L
SEQ ID 8
       EPOPAGROISIRPLLEHCENTHMTIWLGIV
SEO ID 9
SECIDIO E KARGVLKODIS DLS LICS LGYS I LL MVTC
SEO 10 6
       KSRKLPENFNEAKF!TFSMLIFFIVWISF!
SEO ID 7
       GIFYGYKGLLLLGIFLAYE
                                    TKSVSTEKIN
       GIFYGYKGLLLLLGIFLAYETKSVSTEKIN
SEC ID 8
       YAYKGLLMLFGCFLAWETANVSIFALNOSK
SEQ ID 9
SEQ 10:10
       TVYAIKTRGVPETFNEAKPIGFTMYT
```

THLE: G-Protein Fusion Receptors And Chimeric Gaba_b Receptors INVENTORS: Thomas STORMANN, et al. SN: 09/679,064 ATTY #: (072827-1801)

```
      SEQID 6
      PAYASTYGKFVSAVEVIAILAASFGLLACI

      SEOID 7
      DHRAVGMAIYNVAVLCLITAPVTMILSSQQ

      SEOID 8
      DHRAVGMAIYNVAVLCLITAPVTMILSSQQ

      SEOID 9
      YIGMSVYNVGIMCIIGAAVSFLTRDQPNVQ

      SEOID 10
      WLAFIPIFFGTAQSAEKMYIQTTTLTVSMS

      SEOID 7
      DAAFAFASLAIVFSSYITLVVLFVPKM

      SEOID 8
      DAAFAFASLAIVFSSYITLVVLFVPKM

      SEOID 9
      FCIVALVIIFCSTITLCLVFVPKM

      SEOID 9
      FCIVALVIIFCSTITLCLVFVPKL

      SEOID 10
      LSASVSLGMLYMPKVYIIIF
```

```
SECID 11 K P S R N T LE E V R C S T A A H A F K V A A R A T L R R S
SECID 12 R R L L T R G E W Q S E A C D T M K T G S S T N N N E E E K
SEQID 13 R P L I TR G E W C S E A C D T M K T G S S T N A N E E E K
SEQID 14 I T L R T N P D A A T Q N R R F C F T Q N C K K E D S K T S
SEO 10 15 HPEQNVCKAKAS FKAVVTAATMOSKLICKG
SECIDII NVS PKRSSSLGGSTGSTPSSSISSKSNSED
SECIDI2 SRLLEKENRELEKITAEKEERVSELAHOLO
SEO IO 13 S A L L E K E N R E L E K I I A E K E E A V S E L R H C L C
SEQ 10 14 TSVTSVNQASTSRLEGLOSENHRLAMKITE
SEQID 15 NDRPNGEVKSELCESLETNSKSSVEFPMVK
SEO 10 11 P F P Q P E R Q K Q Q C P L A L T Q Q E C Q Q P L T L P Q
SEQID 12 S R Q Q L R S R R H P P T P P E P S G G L P R G P P E P P D
SEOID 13 S R Q Q L R S R R H P P T P P E P S G G L P R G P P E P P D
SECID 14 LOKOLEEV TMQLQD TP EKTTY I KONHYQEL
SEQID 15 SGSTS
SEQID 11 QQR SQQQP R C K Q K V I F G S G T V T F S L S F D E P
SEQID 12 RLSCOGSRVHLLYK
SEQID 13 R L S C D G S R V H L L Y K
SEQID 14 ND ! LNLGNFTESTOGGKA! LKNHLDQNPCL
SEQ 10 15
SEQIDII OKNAMAHGNSTHONSLEADKSSOTLTRHOP
SEQ ID 12
SEQ ID 13
SEQID 14 QWNTTEPSRTCKDPIED INSPEHIQRRLSL
SEQ ID 15
SEQ1011 LLPLQCGETDLOLTVGETGLQGPVGGDGRP
SEQ ID 12
SEQ ID 13
SEQID 14 Q L P I L H H A Y L P S I G G V D A S C V S P C V S P T A S
SEQ 10 15
SEQID 1: EVED PEELS PALV VSSS QS FV ISG GG S T V T
SEO ID 12
SEQ ID 13
SEOID 14 PRHRHVPPS FRVMVSGL
5EQ ID 15
```

Figure 3a

FITE: G-Protein Fusion Receptors And Chimeric Gaba, Receptors INVENTORS: Thomas STORMANN, et al. SN: 09/679,664 ATTY #: (072827-1801)

SEGID 11 ENVVNS

SEQ ID 12

SEQ ID 13

SEQ ID 14

SEQ ID 15

HTLE: G-Protein Fusion Receptors And Chimeric Gaba, Receptors INVENTORS: Thomas STORMANN, et al. SN: 09/679,664 ATTY #: (072827-1801)

SEQ. ID. NO. 16 SEQ. ID. NO. 17	M A E S M A E S	L TWGC L TWRC	C P W C L	TEEEKT TEOEKA	A A R I D C E I N R A A R V D C E I N R
SEQ. ID. NO. 16 SEQ. ID. NO. 17	1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	оккор оккое	REELK	<	GESGKSTFIK GESGKSTFIK
SEQ. ID. NO. 16 SEQ. ID. NO. 17	O M R I O M R I	l H G V G l H G A G	YSEEC YSEEE	D R R A F R L E P K G F R P	L I Y Q N I F V S M L V Y Q N I F V S M
SEC. ID. NO. 16 SEO. ID. NO. 17	C A M I : R A M I I	D A M D R E A M E R	LQIPE	F S R P D S K F S R P E S K	Q H A S L V M T G D H H A S L V M S G D
SEO, ID. NO. 16 SEQ. ID. NO. 17	P Y K V	S T F E K T T F E K	PYAVA PYAAA	R W L W D M A	D A G I R A C Y E R D A G I R A C Y E R
					S E D S Y I P T A Q T E E G Y V P T A Q
					KTKLR I V D V G KTNLR I V D V G
					Y L A S L S E Y D Q Y L A S L S E Y D Q
					I L E L P W F K S T I L E L P W F K S T
					LATYFPSFÖG LATYFPSFÖG
					A
					S V R S V F K D V R N I R K V F K D V R

Figure 4a

TITLE: G-Protein Fusion Receptors And Chimeric Gaba_b Receptors INVENTORS: Thomas STORMANN, et al. SN: 09/679,664 ATTY #: (972827-1801)

SEQ. ID. NO. 16 C S V L A R Y L D E I N L L SEQ. ID. NO. 17 D S V L A R Y L D E I N L L

Figure 4b

ClustalW Formatted Alignments

SEQ. SEQ. SEQ. SEQ.	ID. ID.	NO. NO.	19 20	A A	T T	G G	T G	T G	G G	C C	T C	G C	T C G C	T G	G G	C G	T C	G C	C C	T C	A T	C T	T T	G T	G G	C C	G C	C
SEQ. SEQ. SEQ. SEQ.	ID. ID.	NO. NO.	19 20	C G	A G	C G	T T	C G	T G	T G	C G	C T	C T G C	C G	C C	G C	C A	C C	C T	C G	C C	C C	G G	G C	G T	C T	G C	C T
SEQ. SEQ. SEQ. SEQ.	ID. ID.	NÖ. NO.	19 20	G G	G G	G T	C T	G G	G T	G G	G A	C T	C	C G	A C	G G	A G	C C	C A	C G	C G	Ċ	A G	A T	C G	G G	C	C T
SEQ. SEQ. SEQ. SEQ.	ID. ID.	NO. NO.	19 20	A C	C C	C G	T G	C T	A G	G T	A G	A G	A G G C	G C	T C	T T	G C	C C	C C	A A	G C	A T	T C	C C	A C	T C	A C	C C
SEQ. SEQ. SEQ. SEQ.	ID. ID.	NO. NO.	19 20	A A	C T	C C	C T	G C	C C	C C	C	T C	Т G С	G G	G C	A C	A T	G C	G A	G C	G T	G C	C	A C	T G	C G	A G	G T
SEQ. SEQ. SEQ. SEQ.	ID. ID.	NO. NO.	19 20	G C	T C	A C	C	C C	G C	G G	G C	G A	A C C	C C	Т С	G C	A T	C C	T C	C T	G C	G A	G G	A A	C A	C C	A G	G G
SEQ. SEQ. SEQ.	ID. ID.	NO. NO.	19 20	G C	T G	G C	A G	A C	G A	G G	C T	T G	A A T C	T A	C C	A A	A T	C C	T G	T G	C G	C G	T C	G A	C C	C T	A G	G T
SEQ. SEQ. SEQ.	ID. ID.	NO. NO.	19 20	T T	G T	G C	A C	C C	T A	A T	T G	G A	G A G T	G C	A G	T G	T G	G G	A G	G C	T T	A G	T G	G C	T C	G A	T G	G G

TITLE: G-Protein Fusion Receptors And Chimeric Gaba, Receptors INVENTORS: Thomas STORMANN, et al. SN: 09/679,664 ATTY #: (072827-1801)

SEQ. SEQ. SEQ. SEQ.	ID. ID.	NO. NO.	19 20	C	C G	G G	G C	G C	G A	G G	G	A C	G C	C T	G G	C	G C	A A	G G G C	G C	T C	G C	G G	T C	G G	G G	G T	G G
SEQ. SEQ. SEQ. SEQ.	ID. ID.	NO.	19 20	C G	C A	C G	A A	A T	G G	G G	T C	C G	C C	G T	C G	A G	A A	G G	A T G G	G A	C C	C G	T T	G G	G A	C A	C T	A A
SEQ. SEQ. SEQ.	ID. ID.	NO.	19 20	A G	C	G C	G G	C C	T A	C G	C G	T G	G A	G C	A A	C T	A C	G C	C A T G	T G	A C	T C	G G	G G	A A	C	A T	C A
SEQ. SEQ. SEQ.	. ID. . ID.	NO.	. 19 . 20	A T	C	C A	C G	A C	G T	C	C A	G A	C G	T C	G T	T C	G A	T T	A C C C	C	G A	A C	A C	T A	C	T G	G A	C C
SEQ SEQ SEQ SEQ	. ID. . ID.	. NO . NO	. 19 . 20	T A	C G	C	A A	A A	G G	T T	C G	T T	T	A A	T T	T C	T C	G A	A A G A	C	C	С	T A	G A	G G	A C	A C	A A
SEQ SEQ SEQ	. ID . ID	. NO . NO	. 19 . 20	A C	T C	G A	G A	G	A T	A A	G	G C	T T	T A	T	T	C	C G	T A T	G G	A C	C T	G G	G	G T	T C	G T	G A
SEQ SEQ SEQ	. ID . ID	. NO . NO	. 19 . 20	G C	G A	A A	C	C	T A	C	C	C	A T	G	C T	C	C	T A	T G G A	G A	A T	C	G A	G	A C	G C	C T	C T
SEQ SEQ SEQ SEQ	. ID . ID	. NO . NO	. 19 . 20	C A	G	G G	G	T	G	G	A	. T	T	G	C	C A	G	G C	T T T	G	T	G G	A T	C	C T	C	C C	G A